RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number:	10156	231	S36	
Source:	1	7		
Date Processed by STIC:		13	06	

ENTERED



PCT

RAW SEQUENCE LISTING DATE: 01/13/2006
PATENT APPLICATION: US/10/563,536 TIME: 09:42:53

Input Set : A:\SEQLIST.txt

Output Set: N:\CRF4\01132006\J563536.raw

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5 <110> APPLICANT: WIDMANN, Christian
             YANG, Jiang-Yang
      6
     7
             MICHOD, David
     11 <120> TITLE OF INVENTION: RasGAP derived peptide for selectively killing cancer cells
     15 <130> FILE REFERENCE: KZI-004US
C--> 18 <140> CURRENT APPLICATION NUMBER: US/10/563,536
C--> 18 <141> CURRENT FILING DATE: 2005-12-28
     18 <150> PRIOR APPLICATION NUMBER: PCT/IB2004/002165
     19 <151> PRIOR FILING DATE: 2004-06-29
     21 <150> PRIOR APPLICATION NUMBER: US 60/483,691
     23 <151> PRIOR FILING DATE: 2003-06-30
     27 <160> NUMBER OF SEQ ID NOS: 17
     31 <170> SOFTWARE: PatentIn version 3.1
     35 <210> SEQ ID NO: 1
     37 <211> LENGTH: 249
     39 <212> TYPE: DNA
     41 <213> ORGANISM: Homo sapiens
     45 <400> SEQUENCE: 1
     46 gaagatagaa ggcgtgtacg agctattcta ccttacacaa aagtaccaga cactgatgaa
     48 ataagtttct taaaaggaga tatgttcatt gttcataatg aattagaaga tggatggatg
                                                                              120
     50 tgggttacaa atttaagaac agatgaacaa ggccttattg ttgaagacct agtagaagag
                                                                              180
     52 gtgggccggg aagaagatcc acatgaagga aaaatatggt tccatgggaa gatttccaaa
                                                                              240
     54 caggaagct
                                                                              249
     57 <210> SEQ ID NO: 2
     59 <211> LENGTH: 204
     61 <212> TYPE: DNA
     63 <213> ORGANISM: Homo sapiens
     67 <400> SEQUENCE: 2
     68 gtacgageta ttetacetta cacaaaagta ecagacaetg atgaaataag tttettaaaa
                                                                               60
     70 qqaqatatqt tcattqttca taatqaatta qaaqatqqat gqatgtgggt tacaaattta
                                                                              120
     72 agaacagatg aacaaggcct tattgttgaa gacctagtag aagaggtggg ccgggaagaa
                                                                              180
                                                                              204
     74 gatccacatg aaggaaaaat atgg
     77 <210> SEQ ID NO: 3
     79 <211> LENGTH: 174
     81 <212> TYPE: DNA
     83 <213> ORGANISM: Homo sapiens
     87 <400> SEQUENCE: 3
     88 gtacgagcta ttctacctta cacaaaagta ccagacactg atgaaataag tttcttaaaa
                                                                               60
     90 ggagatatgt tcattgttca taatgaatta gaagatggat ggatgtgggt tacaaattta
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     92 agaacaqatg aacaaggcct tattqttqaa qacctagtag aagaggtggg ccgg
                                                                              174
     95 <210> SEO ID NO: 4
     97 <211> LENGTH: 30
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99 <212> TYPE: DNA

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101 <213> ORGANISM: Homo sapiens
105 <400> SEQUENCE: 4
106 tggatgtggg ttacaaattt aagaacagat
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109 <210> SEQ ID NO: 5
111 <211> LENGTH: 83
113 <212> TYPE: PRT
115 <213> ORGANISM: Homo sapiens
119 <400> SEQUENCE: 5
121 Glu Asp Arg Arg Arg Val Arg Ala Ile Leu Pro Tyr Thr Lys Val Pro
125 Asp Thr Asp Glu Ile Ser Phe Leu Lys Gly Asp Met Phe Ile Val His
129 Asn Glu Leu Glu Asp Gly Trp Met Trp Val Thr Asn Leu Arg Thr Asp
133 Glu Gln Gly Leu Ile Val Glu Asp Leu Val Glu Glu Val Gly Arg Glu
                            55
137 Glu Asp Pro His Glu Gly Lys Ile Trp Phe His Gly Lys Ile Ser Lys
                        70
138 65
                                            75
141 Gln Glu Ala
145 <210> SEQ ID NO: 6
147 <211> LENGTH: 69
149 <212> TYPE: PRT
151 <213> ORGANISM: Homo sapiens
155 <400> SEQUENCE: 6
157 Arg Val Arg Ala Ile Leu Pro Tyr Thr Lys Val Pro Asp Thr Asp Glu
161 Ile Ser Phe Leu Lys Gly Asp Met Phe Ile Val His Asn Glu Leu Glu
                                    25
165 Asp Gly Trp Met Trp Val Thr Asn Leu Arg Thr Asp Glu Gln Gly Leu
                               40
169 Ile Val Glu Asp Leu Val Glu Glu Val Gly Arg Glu Glu Asp Pro His
170 50
                            55
173 Glu Gly Lys Ile Trp
174 65
177 <210> SEQ ID NO: 7
179 <211> LENGTH: 59
181 <212> TYPE: PRT
183 <213> ORGANISM: Homo sapiens
187 <400> SEQUENCE: 7
189 Arg Val Arg Ala Ile Leu Pro Tyr Thr Lys Val Pro Asp Thr Asp Glu
193 Ile Ser Phe Leu Lys Gly Asp Met Phe Ile Val His Asn Glu Leu Glu
                                    25
197 Asp Gly Trp Met Trp Val Thr Asn Leu Arg Thr Asp Glu Gln Gly Leu
201 Ile Val Glu Asp Leu Val Glu Glu Val Gly Arg
202 50
                            55
205 <210> SEQ ID NO: 8
207 <211> LENGTH: 10
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DATE: 01/13/2006

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TIME: 09:42:53
                    PATENT APPLICATION: US/10/563,536
                    Input Set : A:\SEQLIST.txt
                    Output Set: N:\CRF4\01132006\J563536.raw
    209 <212> TYPE: PRT
    211 <213 > ORGANISM: Homo sapiens
    215 <400> SEQUENCE: 8
    217 Trp Met Trp Val Thr Asn Leu Arg Thr Asp
    221 <210> SEQ ID NO: 9
    223 <211> LENGTH: 10
    225 <212> TYPE: PRT
    227 <213> ORGANISM: Bos taurus
    231 <400> SEQUENCE: 9
    233 Trp Met Trp Val Thr Asn Leu Arg Thr Asp
                         5
    237 <210> SEQ ID NO: 10
    239 <211> LENGTH: 10
    241 <212> TYPE: PRT
    243 <213> ORGANISM: Mus musculus
    247 <400> SEQUENCE: 10
    249 Trp Met Trp Val Thr Asn Leu Arg Thr Asp
                                             10
    253 <210> SEQ ID NO: 11
    255 <211> LENGTH: 10
    257 <212> TYPE: PRT
     259 <213> ORGANISM: Rattus norvegicus
    263 <400> SEQUENCE: 11
    265 Trp Met Trp Val Thr Asn Leu Arg Thr Asp
    269 <210> SEQ ID NO: 12
    271 <211> LENGTH: 10
    273 <212> TYPE: PRT
     275 <213> ORGANISM: Anopheles albimanus
     279 <400> SEQUENCE: 12
    281 Trp Leu Trp Val Thr Ala His Arg Thr Gly
     285 <210> SEQ ID NO: 13
     287 <211> LENGTH: 10
     289 <212> TYPE: PRT
     291 <213> ORGANISM: Drosophila melanogaster
     295 <400> SEQUENCE: 13
     297 Trp Leu Trp Val Thr Ala His Arg Thr Gly
    298 1
    301 <210> SEQ ID NO: 14
     303 <211> LENGTH: 10
     305 <212> TYPE: PRT
     307 <213> ORGANISM: Homo sapiens
     311 <220> FEATURE:
     313 <221> NAME/KEY: MISC FEATURE
     315 <222> LOCATION: (2)..(2)
    317 <223> OTHER INFORMATION: Xaa corresponds to an amino acid residue that can be cahnged
by con
               servative or non-conservative amino-acid substitution.
```

RAW SEQUENCE LISTING

DATE: 01/13/2006

```
PATENT APPLICATION: US/10/563,536
                                                             TIME: 09:42:53
                     Input Set : A:\SEQLIST.txt
                     Output Set: N:\CRF4\01132006\J563536.raw
     322 <220> FEATURE:
     324 <221> NAME/KEY: MISC FEATURE
     326 <222> LOCATION: (6)..(7)
     328 <223> OTHER INFORMATION: Xaa correspond to amino acid residues that can be changed by
conser
               vative or non-conservative amino-acid substitutions.
     329
     333 <220> FEATURE:
     335 <221> NAME/KEY: MISC FEATURE
     337 <222> LOCATION: (10)..(10)
     339 <223> OTHER INFORMATION: Xaa corresponds to an amino acid residue that can be changed
by con
     340
               servative or non-conservative amino-acid substitution.
     343 <220> FEATURE:
     345 <221> NAME/KEY: MISC FEATURE
     347 <222> LOCATION: (10)..(10)
     349 <223> OTHER INFORMATION: Xaa corresponds to an amino acid residue that can be changed
by con
     350
               servative or non-conservative amino-acid substitution.
     354 <400> SEQUENCE: 14
W--> 356 Trp Xaa Trp Val Thr Xaa Xaa Arg Thr Xaa
     357 1
     359 <210> SEQ ID NO: 15
     360 <211> LENGTH: 10
     361 <212> TYPE: PRT
     362 <213> ORGANISM: Artificial Sequence
     364 <220> FEATURE:
     365 <223> OTHER INFORMATION: Synthetic Construct
     367 <400> SEQUENCE: 15
     368 Gly Arg Lys Lys Arg Arg Gln Arg Arg Arg
     369 1
     372 <210> SEQ ID NO: 16
     373 <211> LENGTH: 22
     374 <212> TYPE: PRT
     375 <213> ORGANISM: Artificial Sequence
     377 <220> FEATURE:
     378 <223> OTHER INFORMATION: Synthetic Construct
     380 <400> SEQUENCE: 16
     381 Gly Arg Lys Lys Arg Arg Gln Arg Arg Gly Gly Trp Met Trp Val
     382 1
                          5
     383 Thr Asn Leu Arg Thr Asp
     384
                     20
     387 <210> SEQ ID NO: 17
     388 <211> LENGTH: 12
     389 <212> TYPE: PRT
     390 <213> ORGANISM: Artificial Sequence
     392 <220> FEATURE:
     393 <223> OTHER INFORMATION: Synthetic Construct
     395 <400> SEOUENCE: 17
     396 Met Gly Tyr Pro Tyr Asp Val Pro Asp Tyr Ala Ser
     397 1
                          5
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RAW SEQUENCE LISTING

RAW SEQUENCE LISTING ERROR SUMMARY
PATENT APPLICATION: US/10/563,536

DATE: 01/13/2006 TIME: 09:42:54

Input Set : A:\SEQLIST.txt

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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:14; Xaa Pos. 2,6,7,10

VERIFICATION SUMMARY

DATE: 01/13/2006

PATENT APPLICATION: US/10/563,536

TIME: 09:42:54

Input Set : A:\SEQLIST.txt

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L:18 M:270 C: Current Application Number differs, Replaced Current Application No L:18 M:271 C: Current Filing Date differs, Replaced Current Filing Date L:356 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:14 after pos.:0